IN THE SPECIFICATION

Please replace the "Detailed Description of the Drawing" with the following:

FIG. 1 presents the comparative amino acid sequence of **the following members of** the cystein-rich protein family of growth-regulating proteins**: Cyr61 (SEQ ID NO:2); CEF10 (SEQ ID NO:20); Fisp12 (SEQ ID NO:6); CTGF (SEQ ID NO:8): and Nov (SEQ ID NO:21)**.

Please replace the paragraph beginning at page 23, line 13 of the specification as filed with the following:

Alternatively, human *cyr61* cDNA clones were identified with probes generated by RT-PCR. In particular, the prove for screening the human placental cDNA library was a PCR fragment generated with degenerate primers by RT-PCR of total RNA from logarithmically growing WI38 cells. The primers were derived from the sequences corresponding to the most conserved region of the open reading frame of the mouse cyr61 cDNA. One primer, designated H61-5 [5'GGGAATTCTG(TC)GG(GATC)TG(TC)TG(TC)AA(GA)GT(GC)TG-3' **(SEQ ID NO: 18)**], contains a degenerate sequence, which, with the exception of the "GGGAATTC" sequence at the 5' end which was used to introduce an *Eco*RI site, is derived from nucleotides 327-346 (sense strand) of the mouse *cyr61* sequence set forth in SEQ ID NO:1. The degeneracies appear in positions corresponding to the third position of codons in SEQ ID NO:1. The second primer used for PCR amplification of a human *cyr61* sequence was designated H61-3 [5'-CCGGATCC(GA)CA(GA)TT(GA)TA(GA)TT(GA)CA-3' **(SEQ ID NO: 19)**], which, with the exception of the 5' sequence "CCGGATCC" used to introduce a BamHI site, corresponds to the anti-sense strand complementary to nucleotides 1236-1250 of the mouse *cyr61* sequence set forth in SEQ ID NO:1. The degeneracies occur in positions complementary to the third positions of codons in mouse *cyr61* as set forth in SEQ ID NO:1. The amplified *cyr61* cDNA was cloned into the pBlueScript SK+ vector (Stratagene, La Jolla, CA) and sequenced with a Sequenase II kit (U.S. Biochemicals, Cleveland, OH).